

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

10/597,945

Source:

JFLWD

Date Processed by STIC:

08/24/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebe/efs/downloads/documents.htm>>, EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06



I.F.W.O.

RAW SEQUENCE LISTING DATE: 08/24/2006
PATENT APPLICATION: US/10/597,945 TIME: 13:07:45

Input Set : A:\pto.da.txt
Output Set: N:\CRF4\08232006\J597945.csv

2 <110> APPLICANT: Korea Research Institute of Bioscience and Biotechnology
3 . KWAK, Sang-Soo
4 . KWON, Suk-Yoon
5 . LEE, Haeng-Soon
6 . RYU, Sun-Hwa
7 <120> TITLE OF INVENTION: Multiple stress-inducible peroxidase promoter derived from
8 Ipomoea batatas
9
10 <130> FILE REFERENCE: 58049-00035
11 <140> CURRENT APPLICATION NUMBER: US 10/597,945
12 <140> 7006-38-11 > Pb Insert Manditory C1417
13 <141> CURRENT FILING DATE: 2006-08-11
14 <150> PRIOR APPLICATION NUMBER: PCT/KR2004/000381
15 <150> 2004-02-24 > Pb Insert Manditory C1517
16 <170> SOFTWARE: Kopatentplus 1.71
17 <180> NUMBER OF SEQ ID NOS: 30

Does Not Comply
Corrected Diskette Needed

(pg-2,3,4)

ERRONEOUS SEQUENCES

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/597,945

DATE: 08/24/2006

TIME: 13:07:45

Input Set : A:\pto.dn.txt

Output Set: N:\CRF4\08232006\J597945.raw

67	cggacaagg	cttctgttct	cggacaacgc	gtcgatgaat	agcaacaccc	ccagcggcgc	1200
69	cgtaggagg	tacaggagc	agatggacgc	tttttcgcgc	gattttcgccg	ccggccatgg	1260
71	aaagatgagc	cttcctggcc	cgtccccccgg	agtggggctc	gaaatcccgag	aggtttgccg	1320
73	cgagggtgaat	gccaacacag	tttgcattcat	gtggaaatcc	ttccatoga	catacaataac	1380
75	gtctgttatt	ctgtgaaatg	tttactccgg	ctgtgaaagaa	ttttaacttt	ctgtttttt	1440
77	tgaatataaaa	aaagatttttt	ttttatgtcc	taacaaaaact	tgttattactg	aataaaaaattt	1500

E--> 79 ataaatttgt ta

80 1512

1512

119 <210> SEQ ID NO: 4
 120 <211> LENGTH: 306
 121 <212> TYPE: DNA
 122 <213> ORGANISM: Ipomoea batatas
 124 <220> FEATURE:
 125 <221> NAME/KEY: promoter
 126 <222> LOCATION: (1)..(306)
 127 <223> OTHER INFORMATION: -306 deletion promoter

130 <400> SEQUENCE: 4

131	taagggtgtt	tatctgttgc	gcatggatgc	atgacaacacg	catatattat	tataaaaca	60
133	aaatragactt	ccaaat	ttttat	ttttat	ttttat	ttttat	120
135	aaatggaaaca	aaataaaatct	cgttttgcct	tatattat	ttatcaacaa	taataat	180
137	atactgtatcg	aagaatcttc	catttcacgt	tctttat	ttttat	aggaaagctg	240
139	aatccatcatc	atcagcttgc	caacttcattt	ttttat	ttttat	agaagccatt	300

E--> 141 atatcc

142 306

306

145 <210> SEQ ID NO: 5
 146 <211> LENGTH: 366
 147 <212> TYPE: DNA
 148 <213> ORGANISM: Ipomoea batatas
 150 <220> FEATURE:
 151 <221> NAME/KEY: promoter
 152 <222> LOCATION: (1)..(366)
 153 <223> OTHER INFORMATION: -366 deletion promoter

156 <400> SEQUENCE: 5

157	tagtatata	aaataaaatgt	aatccatttc	tatatttgtt	gtatgttattt	agtatcatgg	60
159	taatgggttt	tatctgttgc	gcatggatgc	atgacaacacg	catatattat	tataaaaca	120
161	aaatagtact	ccaaat	ttttat	ttttat	ttttat	ttttat	180
163	aaatggaaaca	aaataaaatct	cgttttgcct	tatattat	ttatcaacaa	taataat	240
165	atactgtatcg	aagaatcttc	catttcacgt	tctttat	ttttat	aggaaagctg	300
167	aatccatcatc	atcagcttgc	caacttcattt	ttttat	ttttat	agaagccatt	360

E--> 169 atatcc

170 366

366

173 <210> SEQ ID NO: 6
 174 <211> LENGTH: 433
 175 <212> TYPE: DNA
 176 <213> ORGANISM: Ipomoea batatas
 178 <220> FEATURE:
 179 <221> NAME/KEY: promoter
 180 <222> LOCATION: (1)..(433)
 181 <223> OTHER INFORMATION: -433 deletion promoter

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/597,945

DATE: 08/24/2006
TIME: 13:07:45

Input Set : A:\pto.ds.txt
Output Set: N:\CRF4\08232006\J597945.raw

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184 <400> SEQUENCE: 6
185 atgtggatct aaaggcgtga atccaaacata tatttcgtaca tttaaaaatg ctaacgtacg 60
187 gttagattag tataatggaa taagttaat catttcttat atttgatgtat ggttaatagt
189 atcaatggtaa ggtttttttt ctgtgcgcga tgatgtcgat acaaaccatg atattttat 180
191 taaaacaaaaa tagacttca atccataaa attatcttat attatartgc caacaatata 240
193 aaatctcaaa tagaaacaat taatcttcgtat tttgttttat attatattat tcaacatata 300
195 taatdtataa ctgcgcgaa aactttttt ttcaactgtt ctatthaagg aagctcgaga 360
197 agccatataa cttccatatec agctcgacca ctccatctt cttccatactt cttttgtgt 420
199 gatatactate atc
200 433
360 <210> SEQ ID NO: 10
361 <211> LENGTH: 1934
362 <212> TYPE: DNA
363 <213> ORGANISM: Ipomea batatas
365 <220> FEATURE:
366 <221> NAME/KRY: promoter
367 <222> LOCATION: (1)..(1934)
368 <223> OTHER INFORMATION: -1934 deletion promoter
371 <40> SEQUENCE: 10
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372-ttgcataatc accatctgtt cttccaaauatc ctaggatattt cttccatactt 60
374 aagtgttccaa ctgttgcataatc atcgaaactt ttacactatc aatttcttccaa tataccctat 120
376 aatataatctt ctactttaaac tagagatttt tccaaactctt: attaatctc ccaaggtaac 180
378 tccccaataa tccaaatggaa aggtttccaaatc ttccaaacttccatccaaacc accggcgtat 240
380 atccataatca tattccatata catataatgtt ttccatctgtt ccctgttccaaatgaaatccatgt 300
382 ttggcgcgtt cggaaaggatc gacggcgtatc caatgttccatccaaactttt tttccatgtt 360
384 attttttatgg tagaaatccc acttataatgtt ttccatgtatcc ataaaaaaatgtt ttggcacaactt 420
386 aggtttccaaatc atttacacacca accaaatccatc ttccatgttccatccaaactttt tttccatgtt 480
388 ttgtttcttc tggaccatgtt ttggcaaaaca atttttgaac caccacttata ctacttccaaa 540
390 attatgttataa ttttttttatgtt agttttttccat cttttagatgc tacatgtatccatccaaatgtt 600
392 gttccaaatataa ctttccatgtt ttccatccaaatc tttttccatccaaatgtt ccatttccatccaaatgtt 660
394 cttgtttttttt ctttccatgtt ttccatccaaatc ataaatgtt atttttttccatccaaatgtt 720
396 aatggatcatgtt accaaatccatc ggtttttccatccaaatgtt ccatttccatccaaatgtt 780
398 taatgtatataa gtataatctt ttgtaaaaat ttgtttttttt tttttgtatccatccaaatgtt 840
400 ttccaaatgtt ttccatccatccaaatgtt ttgtttttttt tttttgtatccatccaaatgtt 900
402 atttttttccatccaaatgtt ttgtttttttt tttttgtatccatccaaatgtt 960
404 aaacaaatataa accaaatataa tggaaatggta atttttttccatccaaatgtt ttgtttttttt 1020
406 ctgtatccatccaaatgtt ttgtttttccatccaaatgtt ttgtttttttt tttttgtatccatccaaatgtt 1080
408 gtataatataa actttttttccatccaaatgtt ttgtttttccatccaaatgtt ttgtttttttt tttttgtatccatccaaatgtt 1140
410 ttccatccatccaaatgtt ttgtttttccatccaaatgtt ttgtttttttt tttttgtatccatccaaatgtt 1200
412 gatattttatccatccaaatgtt ttgtttttccatccaaatgtt ttgtttttttt tttttgtatccatccaaatgtt 1260
414 daatccatccaaatgtt ttgtttttccatccaaatgtt ttgtttttttt tttttgtatccatccaaatgtt 1320
416 ggttactatccatccaaatgtt ttgtttttccatccaaatgtt ttgtttttttt tttttgtatccatccaaatgtt 1380
418 atttatccatccaaatgtt ttgtttttccatccaaatgtt ttgtttttttt tttttgtatccatccaaatgtt 1440
420 atttatccatccaaatgtt ttgtttttccatccaaatgtt ttgtttttttt tttttgtatccatccaaatgtt 1500
422 aatggtgactt taatggcgtatc aatccaaatccatccaaatgtt ttgtttttccatccaaatgtt 1560
424 ggtttagatccatccaaatgtt ttgtttttccatccaaatgtt ttgtttttttt tttttgtatccatccaaatgtt 1620
426 tatccatgtttaa aggtttttt ccattttccatccaaatgtt ttgtttttccatccaaatgtt 1680
428 tttttttccatccaaatgtt ttgtttttccatccaaatgtt ttgtttttttt tttttgtatccatccaaatgtt 1740
430 aaatccatccaaatgtt ttgtttttccatccaaatgtt ttgtttttttt tttttgtatccatccaaatgtt 1800

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/597,945

DATE: 08/24/2006

TIME: 13:07:45

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\08232006\J597945.raw

432 ataatttaat actgatecgaa gaactttccc ttcaagttc tctatattaag gaaggctttag 1860
434 aagccattaa tcctccatcat cagtcgacc acctattttct tccttcatact tccttttgtg 1920
E--> 435 tgataatcat catc
437 1934
770 <210> SEQ ID NO: 30
771 <211> LENGTH: 21
772 <212> TYPE: DNA
773 <213> ORGANISM: Artificial Sequence
775 <220> FEATURE:
776 <223> OTHER INFORMATION: reverse primer for NPTII
779 <400> SEQUENCE: 30
780 atccgggacgc gogataccgt a

E--> 782 1
E--> 786 1

1934

21

delete

RAM SEQUENCE LISTING ERROR SUMMARY DATE: 08/24/2006
PATENT APPLICATION: US/10/597,945 TIME: 13:07:46

Input Set : A:\pto.da.txt
Output Set: N:\CRF4\08232006\J597945.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:4; Line(s) 141
Seq#:5; Line(s) 169

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/597,945

DATE: 08/24/2006

TIME: 13:07:46

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\08232006\J597945.raw

L:14 M:259 W: Allowed number of lines exceeded, <140> CURRENT APPLICATION NUMBER:
L:16 M:271 C: Current Filing Date differs. Replaced Current Filing Date
L:17 M:259 W: Allowed number of lines exceeded, <150> PRIOR APPLICATION NUMBER:
L:79 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:1512 SEQ:1
L:141 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:306 SEQ:4
L:169 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:366 SEQ:5
L:199 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:433 SEQ:6
L:436 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:1934 SEQ:10
L:783 M:254 E: No. of Bases conflict, this line has no nucleotides.
L:786 M:254 E: No. of Bases conflict, this line has no nucleotides.